## **AMENDMENTS TO THE CLAIMS**

Claim 1. (Currently amended) A formulation of thermostable DNA polymerase comprising at least one thermostable DNA polymerase lacking 3'-exonuclease activity and at least one thermostable DNA polymerase exhibiting 3'-exonuclease activity wherein the ratio of DNA polymerase activity of the at least one polymerase lacking 3'-exonuclease activity to DNA polymerase activity of the at least one polymerase exhibiting 3'-exonuclease is greater than 1 to 1. from about 100:1 up to about 600:1.

Claim 2. (Currently amended) A formulation of thermostable DNA polymerase comprising at least one thermostable DNA polymerase lacking 3'-exonuclease activity and at least one thermostable DNA polymerase exhibiting 3'-exonuclease activity wherein the ratio of DNA polymerase activity of the at least one polymerase lacking 3'-exonuclease activity to DNA polymerase activity of the at least one polymerase exhibiting 3'-exonuclease activity is greater than 1 to 1 and wherein the at least one thermostable DNA polymerase lacking 3'-exonuclease activity is selected from the group consisting of Klentaq-291 and Klentaq-278.

Claim 3. (Cancelled)

Claim 4. (Currently amended) A formulation of DNA polymerase <u>as set</u> <u>forth in claim 1</u> comprising at least one DNA polymerase which lacks 3'-exenuclease activity, and at least one DNA polymerase which exhibits 3'-exenuclease activity wherein the at least one DNA polymerase which exhibits 3'-exenuclease activity is selected from the group consisting of an Archaebacterial DNA polymerase and wherein the ratio of DNA polymerase activity of the at least one polymerase which lacks 3' exenuclease activity to DNA polymerase activity of the at least 4:1.

Claim 5. (Previously presented) A formulation of DNA polymerase as set forth in claim 4 wherein the Archaebacterial DNA polymerase is selected from the group consisting of a Pyrococcus furiosus DNA polymerase, a Thermococcus litoralis DNA polymerase, and a combination thereof.

Claim 6. (Previously presented) A formulation of DNA polymerase as set forth in claim 5 wherein the Archaebacterial DNA polymerase is a Pyrococcus furiosus DNA polymerase and wherein the ratio of DNA polymerase activity of the at least one polymerase which lacks 3'-exonuclease activity to DNA polymerase activity of the at least one polymerase which exhibits 3'-exonuclease activity is from about 150:1 to about 170:1.

## Claims 7-9. (Canceled)

Claim 10. (Currently amended) A formulation of DNA polymerase as set forth in claim 4 wherein the at least one polymerase which lacks 3'-exonuclease activity is selected from the group consisting of a wild-type Thermus aquaticus DNA polymerase and a mutein of a Thermus aquaticus DNA polymerase from which the N-terminal 3 amino acids have been deleted, wherein the at least one polymerase which lacks 3'-exonuclease activity is a Pyrococcus furiosus DNA polymerase, and wherein the ratio of DNA polymerase activity of the at least one polymerase which lacks 3'-exonuclease activity to DNA polymerase activity of the at least one polymerase which exhibits 3'-exonuclease activity is from about 10:1 to about 15:1.

Claim 11. (Previously presented) A formulation of DNA polymerase as set forth in claim 5 wherein the at least one Archaebacterial DNA polymerase which exhibits 3'-exonuclease activity is selected from the group consisting of a Thermus flavus DNA polymerase and a Thermus thermophilus DNA polymerase.

## Claims 12-16. (Cancelled)

Claim 17. (Currently amended) A method as set forth in claim 14 for amplifying a nucleic acid, comprising:

preparing a composition comprising a DNA polymerase comprising at least one thermostable DNA polymerase which lacks 3'-exonuclease activity, at least one thermostable DNA polymerase which exhibits 3'-exonuclease activity, and a nucleic acid comprising a sequence to be amplified, wherein the ratio of DNA polymerase activity of the at least one thermostable DNA polymerase which lack 3'-exonuclease activity to DNA polymerase activity of the at least one thermostable DNA polymerase which exhibits 3'-exonuclease activity from about 100:1 up to about 600:1; and

wherein the nucleic acid sequence to be amplified is a nucleic acid sequence 6 kb or more in length[ . ]

subjecting the composition to conditions effective for amplifying the nucleic acid sequence.

Claim 18. (Currently amended) A method as set forth in claim 44 17 wherein the nucleic acid sequence to be amplified is a nucleic acid sequence 8.4 kb or more in length.

Claim 19. (Currently amended) A method as set forth in claim 44 <u>17</u> wherein the nucleic acid sequence to be amplified is a nucleic acid sequence 15 kb or more in length.

Claim 20. (Currently amended) A method as set forth in claim 44 <u>17</u> wherein the composition further comprises an oligonucleotide primer or primers, wherein at least one primer is itself a product of a PCR amplification.

Claim 21. (Currently amended) A method as set forth in claim 14 for amplifying a nucleic acid, comprising:

preparing a composition comprising a DNA polymerase comprising at least one thermostable DNA polymerase which lacks 3'-exonuclease activity, at least one thermostable DNA polymerase which exhibits 3'-exonuclease activity, and a nucleic acid comprising a sequence to be amplified, wherein the ratio of DNA polymerase activity of the at least one thermostable DNA polymerase which lack 3'-exonuclease activity to DNA polymerase activity of the at least one thermostable DNA polymerase which exhibits 3'-exonuclease activity exceeds 1 to 1; and wherein the nucleic acid sequence to be amplified is a nucleic acid sequence 6 kb or more in length wherein the conditions effective for amplifying the nucleic acid sequence comprise conditions effective for denaturing the nucleic acid sequence, and wherein the denaturing has a duration of less than 20 seconds

Claim 22. (Previously presented) A method as set forth in claim 21 wherein the denaturing has a duration of less than 5 seconds.

Claims 23-24 (Cancelled)

Claim 25. (Previously presented) A formulation of DNA polymerase as set forth in claim 4 wherein the at least one polymerase which lacks 3'-exonuclease activity is a Thermus thermophilus DNA polymerase.

Claim 26. (Previously presented) A formulation of thermostable DNA polymerase as set forth in claim 1 wherein the at least one thermostable DNA polymerase lacking 3'-exonuclease activity comprises at least one\_thermostable DNA polymerase which in wild-type form lacks any 3'-exonuclease activity.

Claims 27-30. (Cancelled)

Claim 31. (Previously presented) A formulation of DNA polymerase in accordance with claim 5, wherein the at least one polymerase which lacks 3'-exonuclease activity is selected from the group consisting of a 3'-exonuclease-negative mutant form of a DNA polymerase which exhibits 3'-exonuclease activity, a Thermus aquaticus DNA polymerase, a Thermus flavus DNA polymerase, a Thermus thermophilus DNA polymerase and a combination thereof.

Claim 32. (Currently amended) A composition comprising at least one DNA thermostable polymerase lacking 3'-exonuclease activity and at least one thermostable DNA polymerase exhibiting 3-exonuclease activity, wherein the ratio of the at least one DNA polymerase lacking 3'-exonuclease activity to the at least one polymerase exhibiting 3'-exonuclease activity is greater-than-1:1 from about 100:1 up to about 600:1 by weight.

Claim 33. (Currently amended) A composition in accordance with claim 32, wherein the ratio of the at least one DNA polymerase lacking 3'-exonuclease activity to the at least one polymerase exhibiting 3'-exonuclease activity is at least about 4:1 1:150 to about 1:170 by weight.